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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2010; month=1; day=20; hr=13; min=10; sec=55; ms=609;]

=====

Reviewer Comments:

<210> 1

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> sequence for HIS-tag

<400> 1

His His His His His

1 5

As an explanation of "<213> Artificial Sequence": in the above <223> response ("sequence for HIS-tag"), please give more information regarding the source of the genetic material. Please ensure that all subsequent sequences showing "<213> Artificial Sequence" show the source of the genetic material in the <223> response. Please do not show TAB codes in the sequence listing.

<210> 4

<211> 8

<212> PRT

<213> Artificial Sequence

<223> sequence for HIS-tag

<220>

<223> linkage to a monomeric titin I28 Ig domain

<400> 4

Tyr Gly His His His His His His

1 5

Please insert a mandatory <220> above the "<223> sequence for HIS-tag". "<220>" is a mandatory header whenever <221>, <222>, or <223> is shown. Also, please give more information regarding the source of the genetic material of "<223> sequence for HIS-tag".

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<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> sequence for HIS-tag

<220>

<223> linkage to a monomeric titin I28 Ig domain

<400> 5

Tyr Gly Tyr Gly His His His His His His

1 5 10

Besides an insufficient explanation of "<213> Artificial Sequence" (<223> sequence for HIS-tag"), the amino acid numbers in the above sequence are misaligned. Do not use TAB codes between amino acid numbers; TAB's cause misalignment. Do not use TAB codes in the sequence listing.

<210> 9

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> sequence for HIS-tag

<400> 9

His His His Gly Tyr Gly His His His

1 5

Page 4

In the above Sequence 9, please give more information in the <223> response regarding the source of the genetic material.

Please remove "Page 4" at the end of the submitted file.

Application No: 10567992 Version No: 2.0

Input Set:

Output Set:

Started: 2009-12-30 19:29:41.712
Finished: 2009-12-30 19:29:42.990
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 278 ms
Total Warnings: 11
Total Errors: 1
No. of SeqIDs Defined: 9
Actual SeqID Count: 9

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 333	tabs used in amino acid numbering SEQID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 112	Upper case found in data; Found at position(0) SeqId(9)
E 259	Found undefined lettercode; POS (4) SEQID(9)

SEQUENCE LISTING

<110> Stewart, Russell J
Kiser, Patrick F
Staynor, Richard S

<120> Crosslinking Within Coordination Complexes

<130> 007180-50 US

<140> 10567992
<141> 2009-12-30

<150> US 60/494,349
<151> 2003-08-11

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His His His His His
1 5

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<223> glutathione S-transferase at C-terminus

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His His His Gly Tyr Gly His His His

1 5

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